

SEQUENCE LISTING

<110> Rondon, Isaac J
Ladner, Robert C

<120> BINDING PEPTIDES FOR CARCINOEMBRYONIC ANTIGEN (CEA)

<130> Sequence Listing DYX-016.0 US

<140> (not yet assigned)

<141> 2000-04-03

<160> 107

<170> PatentIn Ver. 2.1

<210> 1

<211> 16

<212> PRT

<213> Artificial Sequence

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<223> Description of Artificial Sequence: CEA binding
polypeptide

<220>

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<222> (1)

<223> Xaa is Asn, Asp or is absent

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<221> VARIANT

<222> (2)

<223> Xaa is Trp

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<221> VARIANT

<222> (3)

<223> Xaa is Asp, Phe or Val

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<221> VARIANT

<222> (5)

<223> Xaa is Asn, Glu or Met

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<221> VARIANT

<222> (6)

<223> Xaa is Asn, Leu, Met or Phe

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<223> Xaa is Asp, Gly, Ile, Lys, Phe or Thr

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<223> Xaa is Ala, Gln, gly Lys or Thr

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<221> VARIANT

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<223> Xaa is Arg, Asn, Asp, Glu or Gly

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<223> Xaa is Gln, Leu or Gly

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<223> Xaa is Ala, Trp or Tyr

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<222> (15)

<223> Xaa is Arg, Leu Pro or Ser

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<222> (16)

<223> Xaa is Leu, Ser, Trp or Tyr

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Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa

1

5

10

15

<210> 2

<211> 16

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<223> Description of Artificial Sequence: family of
preferred CEA binding moieties

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<223> Xaa is Arg, Leu, Pro or Ser

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<222> (16)

<223> Xaa is Leu, Ser, Trp or Tyr

<400> 2

Xaa Trp Val Cys Glu Xaa Xaa Lys Xaa Gln Trp Xaa Cys Asn Xaa Xaa
1 5 10 15

<210> 3

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<212> PRT

<213> Artificial Sequence

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<223> Description of Artificial Sequence: CEA binding
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<222> (5)

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<222> (9)
<223> Xaa is Ala, Gly, His, Phe, Thr or Val

<400> 3
Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys
1 5 10

<210> 4
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<220>
<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 4
Asn Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asn Ser Tyr
1 5 10 15

<210> 5
<211> 16
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 5
Asp Trp Val Cys Glu Asn Lys Lys Asp Gln Trp Thr Cys Asn Leu Leu
1 5 10 15

<210> 6
<211> 16
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<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 6

Asn Trp Asp Cys Met Phe Gly Ala Glu Gly Trp Ala Cys Ser Pro Trp
1 5 10 15

<210> 7

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 7

Asp Trp Val Cys Glu Lys Thr Thr Gly Gly Tyr Val Cys Gln Pro Leu
1 5 10 15

<210> 8

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 8

Asn Trp Phe Cys Glu Met Ile Gly Arg Gln Trp Gly Cys Val Pro Ser
1 5 10 15

<210> 9

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 9

Asp Trp Val Cys Asn Phe Asp Gln Gly Leu Ala His Cys Phe Pro Ser
1 5 10 15

<210> 10

<211> 12
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<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: parental
domain for design of microprotein display library

<220>
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<222> (1)..(12)
<223> amino acid positions 4 and 9 are invariant Cys;
all other positions Xaa are varied but not Cys, to
provide a library of 2×10^8 different peptides
based on the template sequence

<400> 10
Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa
1 5 10

<210> 11
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<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: parental
domain for design of microprotein display library

<220>
<221> VARIANT
<222> (1)..(11)
<223> amino acid positions 3 and 9 are invariant Cys;
all other positions Xaa are varied but not Cys, to
provide a library of 1×10^9 different peptides
based on the template sequence

<400> 11
Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa
1 5 10

<210> 12
<211> 12
<212> PRT
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: parental
domain for design of microprotein display library

<220>

<221> VARIANT

<222> (1)..(12)

<223> amino acid positions 3 and 10 are invariant Cys;
all other positions Xaa are varied but not Cys, to
provide a library of 1×10^9 different peptides
based on the template sequence

<400> 12

Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa
1 5 10

<210> 13

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: parental
domain for design of microprotein display library

<220>

<221> VARIANT

<222> (1)..(16)

<223> amino acid positions 4 and 13 are invariant Cys;
all other positions Xaa are varied but not Cys, to
provide a library of 2.5×10^8 different peptides
based on the template sequence

<400> 13

Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa
1 5 10 15

<210> 14

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: variable
sublibrary sequence used in designing focused
secondary library

<220>

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<222> (1)..(3)

<223> Xaa is any amino acid except Cys

<220>

<221> VARIANT

<222> (5)..(6)

<223> Xaa is any amino acid except Cys

<400> 14

Xaa	Xaa	Xaa	Cys	Xaa	Xaa	Lys	Lys	Asp	Gln	Trp	Thr	Cys	Asn	Leu	Leu
1				5					10					15	

<210> 15

<211> 16

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<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: variable
sublibrary sequence used in designing focused
secondary library

<220>

<221> VARIANT

<222> (5)..(9)

<223> Xaa is any amino acid except Cys

<400> 15

Asp	Trp	Val	Cys	Xaa	Xaa	Xaa	Xaa	Xaa	Gln	Trp	Thr	Cys	Asn	Leu	Leu
1				5					10					15	

<210> 16

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: variable
sublibrary sequence used in designing focused
secondary library

<220>

<221> VARIANT

<222> (8)..(12)

<223> Xaa is any amino acid except Cys

<400> 16

Asp	Trp	Val	Cys	Glu	Asn	Lys	Xaa	Xaa	Xaa	Xaa	Xaa	Cys	Asn	Leu	Leu
1					5				10					15	

<210> 17

<211> 16

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<220>

<223> Description of Artificial Sequence: variable
sublibrary sequence used in designing focused
secondary library

<220>

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<222> (11)..(12)

<223> Xaa is any amino acid except Cys

<220>

<221> VARIANT

<222> (14)..(16)

<223> Xaa is any amino acid except Cys

<400> 17

Asp	Trp	Val	Cys	Glu	Asn	Lys	Lys	Asp	Gln	Xaa	Xaa	Cys	Xaa	Xaa	Xaa
1					5				10					15	

<210> 18

<211> 16

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<213> Artificial Sequence

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sublibrary sequence used in designing focused
secondary library

<220>

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<222> (6)..(7)

<223> Xaa is any amino acid except Cys

<220>
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<220>
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<223> Xaa is any amino acid except Cys

<220>
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<222> (15)
<223> Xaa is any amino acid except Cys

<400> 18
Asp Trp Val Cys Glu Xaa Xaa Lys Xaa Gln Trp Xaa Cys Asn Xaa Leu
1 5 10 15

<210> 19
<211> 16
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secondary library

<220>
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<223> Xaa is any amino acid except Cys

<220>
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<222> (9)
<223> Xaa is any amino acid except Cys

<220>
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<222> (12)
<223> Xaa is any amino acid except Cys

<400> 19
Asn Trp Val Cys Xaa Xaa Xaa Lys Xaa Gln Trp Xaa Cys Asn Ser Tyr
1 5 10 15

<210> 20
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<220>
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sublibrary sequence used in designing focused
secondary library

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<223> Xaa is any amino acid except Cys

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<223> Xaa is any amino acid except Cys

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<223> Xaa is any amino acid except Cys

<400> 20
Xaa Trp Xaa Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Xaa Xaa Xaa
1 5 10 15

<210> 21
<211> 16
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: isolate of
TN10/9 library found not to bind CEA

<400> 21
Asn Trp Arg Cys Lys Leu Phe Pro Arg Tyr Pro Tyr Cys Ser Ser Trp
1 5 10 15

<210> 22
<211> 15

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: isolate of
TN10/9 library found not to bind CEA

<400> 22

Arg	Tyr	Cys	Glu	Phe	Phe	Pro	Trp	Ser	Leu	His	Cys	Gly	Arg	Pro
1					5				10					15

<210> 23

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: conserved
amino acid positions in first family of CEA
binding peptides

<220>

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<222> (6)

<223> X is Asn, Leu, Met or Phe

<220>

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<222> (7)

<223> X is Asp, Gly, Ile, Lys, Phe or Thr

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<221> VARIANT

<222> (9)

<223> X is Arg, Asn, Asp, Glu or Gly

<220>

<221> VARIANT

<222> (12)

<223> X is Ala, Gly, His, Phe, Thr or Val

<220>

<221> VARIANT

<222> (15)

<223> X is Arg, Leu, Pro or Ser

<400> 23

Asp	Trp	Val	Cys	Glu	Xaa	Xaa	Lys	Xaa	Gln	Trp	Xaa	Cys	Asn	Xaa	Leu
1				5					10					15	

<210> 24

<211> 27

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: synthetic CEA
binding peptide with C-terminal immobilization
sequence

<400> 24

Ser	Asn	Trp	Val	Cys	Asn	Leu	Phe	Lys	Asn	Gln	Trp	Phe	Cys	Asn	Ser
1				5					10					15	

Tyr	Ala	Pro	Gly	Gly	Glu	Gly	Gly	Gly	Ser	Lys
			20					25		

<210> 25

<211> 27

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: synthetic CEA
binding peptide with C-terminal immobilization
sequence

<400> 25

Ser	Asp	Trp	Val	Cys	Glu	Asn	Lys	Lys	Asp	Gln	Trp	Thr	Cys	Asn	Leu
1				5					10					15	

Leu	Ala	Pro	Gly	Gly	Glu	Gly	Gly	Gly	Ser	Lys
			20					25		

<210> 26

<211> 27

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: synthetic CEA
binding peptide with C-terminal immobilization

sequence

<400> 26

Ser Asn Trp Asp Cys Met Phe Gly Ala Glu Gly Trp Ala Cys Ser Pro
1 5 10 15

Trp Ala Pro Gly Gly Glu Gly Gly Gly Ser Lys
20 25

<210> 27

<211> 27

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: synthetic CEA
binding peptide with C-terminal immobilization
sequence

<400> 27

Ser Asp Trp Val Cys Glu Leu Thr Thr Gly Gly Tyr Val Cys Gln Pro
1 5 10 15

Leu Ala Pro Gly Gly Glu Gly Gly Gly Ser Lys
20 25

<210> 28

<211> 10

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: C-terminal
sequence for immobilizing peptides

<400> 28

Ala Pro Gly Gly Glu Gly Gly Gly Ser Lys
1 5 10

<210> 29

<211> 16

<212> PRT

<213> Artificial Sequence

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<223> Description of Artificial Sequence: template
sequence for sublibrary used in construction of
focused secondary display library

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<222> (1)..(3)

<223> X is any amino acid except Cys

<220>

<221> VARIANT

<222> (5)..(6)

<223> X is any amino acid except Cys

<400> 29

Xaa	Xaa	Xaa	Cys	Xaa	Xaa	Lys	Lys	Asp	Gln	Trp	Thr	Cys	Asn	Leu	Leu
1				5					10					15	

<210> 30

<211> 16

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<223> Description of Artificial Sequence: template
sequence for sublibrary used in construction of
focused secondary display library

<220>

<221> VARIANT

<222> (5)..(9)

<223> X is any amino acid except Cys

<400> 30

Asp	Trp	Val	Cys	Xaa	Xaa	Xaa	Xaa	Xaa	Gln	Trp	Thr	Cys	Asn	Leu	Leu
1				5					10					15	

<210> 31

<211> 16

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<223> Description of Artificial Sequence: template
sequence for sublibrary used in construction of
focused secondary display library

<220>
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 <222> (8)..(12)
 <223> X is any amino acid except Cys

 <400> 31
 Asp Trp Val Cys Glu Asn Lys Xaa Xaa Xaa Xaa Xaa Cys Asn Leu Leu
 1 5 10 15

<210> 32
 <211> 16
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<220>
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 sequence for sublibrary used in construction of
 focused secondary display library

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 <223> X is any amino acid except Cys

<220>
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 <223> X is any amino acid except Cys

<400> 32
 Asp Trp Val Cys Glu Asn Lys Lys Asp Gln Xaa Xaa Cys Xaa Xaa Xaa
 1 5 10 15

<210> 33
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<220>
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 sequence for sublibrary used in construction of
 focused secondary display library

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<222> (6)..(7)

<223> X is any amino acid except Cys

<220>

<221> VARIANT

<222> (9)

<223> X is any amino acid except Cys

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<222> (12)

<223> X is any amino acid except Cys

<220>

<221> VARIANT

<222> (15)

<223> X is any amino acid except Cys

<400> 33

Asp Trp Val Cys Glu Xaa Xaa Lys Xaa Gln Trp Xaa Cys Asn Xaa Leu

1

5

10

15

<210> 34

<211> 16

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<223> Description of Artificial Sequence: template
sequence for sublibrary used in construction of
focused secondary display library

<220>

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<222> (5)..(7)

<223> X is any amino acid except Cys

<220>

<221> VARIANT

<222> (9)

<223> X is any amino acid except Cys

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<221> VARIANT

<222> (12)

<223> X is any amino acid except Cys

<400> 34

Asn Trp Val Cys Xaa Xaa Xaa Lys Xaa Gln Trp Xaa Cys Asn Ser Tyr
1 5 10 15

<210> 35

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: template
sequence for sublibrary used in construction of
focused secondary display library

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<222> (1)

<223> X is any amino acid except Cys

<220>

<221> VARIANT

<222> (3)

<223> X is any amino acid except Cys

<220>

<221> VARIANT

<222> (14)..(16)

<223> X is any amino acid except Cys

<400> 35

Xaa Trp Xaa Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Xaa Xaa Xaa
1 5 10 15

<210> 36

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: family of CEA
binding polypeptides

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<221> VARIANT

<222> (1)

<223> Xaa is Asp, Asn, Ala or Ile

<220>
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 <222> (3)
 <223> Xaa is Val, Ile, Met, Tyr, Phe, Pro or Asp

<220>
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 <223> Xaa is Asn, Glu or Asp

<220>
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<220>
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 <222> (7)
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 Ser, Val, Trp or Tyr

<220>
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 <222> (8)
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<220>
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 <222> (9)
 <223> Xaa is Asn, Pro, Phe, Gly, Asp, Ala, Ser, Glu, Gln
 or Trp

<220>
 <221> VARIANT
 <222> (10)
 <223> Xaa is Gln or Lys

<220>
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 <222> (12)
 <223> Xaa is Phe, Thr, Met, Ser, Ala, Asn, Val, His,
 Ile, Pro, Trp or Tyr

<220>
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 <222> (14)
 <223> Xaa is Asn, Asp, Glu, Pro, Gln or Ser

<220>

<221> VARIANT

<222> (15)

<223> Xaa is Val, Leu, Ile, Pro, Ala, Gln, Ser, Met,
Glu,Thr, Lys or Trp

<220>

<221> VARIANT

<222> (16)

<223> Xaa is Leu, Met, Val, Tyr, Ala, Ile, Trp, His,
Pro, Gln, Glu, Phe, Lys or Arg

<400> 36

Xaa Trp Xaa Cys Xaa Xaa Xaa Xaa Xaa Trp Xaa Cys Xaa Xaa Xaa
1 5 10 15

<210> 37

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 37

Asp Trp Met Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Leu Met
1 5 10 15

<210> 38

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 38

Asp Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Leu Met
1 5 10 15

<210> 39

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 39

Asp	Trp	Ile	Cys	Asn	Leu	Phe	Lys	Asn	Gln	Trp	Phe	Cys	Asp	Gln	Met
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<210> 40

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 40

Asn	Trp	Ile	Cys	Asn	Leu	Phe	Lys	Asn	Gln	Trp	Phe	Cys	Asp	Gln	Glu
1				5					10					15	

<210> 41

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 41

Asp	Trp	Ile	Cys	Asn	Leu	Phe	Lys	Asn	Gln	Trp	Phe	Cys	Gln	Val	Lys
1				5					10					15	

<210> 42

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 42

Asp Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Val Met
1 5 10 15

<210> 43

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 43

Asp Trp Met Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Gln Ile
1 5 10 15

<210> 44

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 44

Ile Trp Asp Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Pro Ala Pro
1 5 10 15

<210> 45

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 45

Asp Trp Ile Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Ile Arg
1 5 10 15

<210> 46
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<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 46

Asp Trp Met Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Val Val
1 5 10 15

<210> 47
<211> 16
<212> PRT
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 47

Asp Trp Ile Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Ala Ile
1 5 10 15

<210> 48
<211> 16
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<400> 48

Asp Trp Ile Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Met Ala
1 5 10 15

<210> 49
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<400> 49

Asp Trp Val Cys Glu Phe Leu Lys Met Gln Trp Ala Cys Asn Val Leu
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<210> 50

<211> 16

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Asp Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asn Val Met
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<210> 51

<211> 16

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<400> 51

Ala Trp Pro Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Pro Pro Gln
1 5 10 15

<210> 52

<211> 16

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<213> Artificial Sequence

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Asp Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Val Leu
1 5 10 15

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1 5 10 15

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Asp Trp Val Cys Glu Trp Leu Lys Met Gln Trp Ala Cys Asn Met Leu
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Asp Trp Val Cys Asp Phe Phe Phe Asn Gln Trp Thr Cys Asn Leu Leu
1 5 10 15

<210> 56
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Asp Trp Val Cys Glu Met Phe Lys Ala Gln Trp Phe Cys Asn Ala Leu
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<400> 57

Asp Trp Ile Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Ala Trp
1 5 10 15

<210> 58

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<400> 58

Asp Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Val Trp
1 5 10 15

<210> 59

<211> 16

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<400> 59

Asp	Trp	Val	Cys	Glu	Tyr	Phe	Lys	Asn	Gln	Trp	Phe	Cys	Asn	Val	Leu
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<210> 60
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Asp	Trp	Val	Cys	Glu	Ile	Asp	Lys	Gly	Gln	Trp	Thr	Cys	Asn	Pro	Leu
1				5					10					15	

<210> 61
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Asp	Trp	Val	Cys	Asn	Leu	Phe	Lys	Asn	Gln	Trp	Phe	Cys	Asn	Pro	Phe
1				5					10					15	

<210> 62
 <211> 16
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Asp	Trp	Val	Cys	Asn	Leu	Phe	Lys	Asn	Gln	Trp	Phe	Cys	Asp	Val	Gln
1				5					10					15	

<210> 63
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<212> PRT
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 63

Asp Trp Val Cys Asn Leu Phe Phe Gly Gln Trp Thr Cys Asn Leu Leu
1 5 10 15

<210> 64

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 64

Asp Trp Ile Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Glu Ala His
1 5 10 15

<210> 65

<211> 16

<212> PRT

<213> Artificial Sequence

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Asp Trp Val Cys Glu Leu Val Lys Ala Gln Trp Tyr Cys Asn Ile Leu
1 5 10 15

<210> 66

<211> 16

<212> PRT

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<400> 66

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1 5 10 15

<210> 67

<211> 16

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Asp Trp Val Cys Glu Phe Tyr Lys Ser Gln Trp Asn Cys Asn Ile Leu
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<210> 68

<211> 16

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Asp Trp Val Cys Glu Trp Phe Lys Pro Gln Trp Phe Cys Asn Pro Leu
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<210> 69

<211> 16

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Asp Trp Tyr Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Val Leu
1 5 10 15

<210> 70
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Asp Trp Val Cys Glu Tyr Asn Asp Glu Gln Trp Thr Cys Asn Leu Leu
1 5 10 15

<210> 71
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<400> 71
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1 5 10 15

<210> 72
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<400> 72
Asp Trp Val Cys Asn Trp Glu Leu Phe Gln Trp Thr Cys Asn Leu Leu
1 5 10 15

<210> 73
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<400> 73

Asp Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Gln Val
1 5 10 15

<210> 74

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<400> 74

Asp Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Val Pro
1 5 10 15

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<400> 75

Asp Trp Val Cys Glu Phe Phe Lys Gln Gln Trp Phe Cys Asn Val Leu
1 5 10 15

<210> 76

<211> 16

<212> PRT

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<400> 76

Asp Trp Val Cys Glu Phe Phe Lys Asp Gln Trp Ser Cys Asn Val Leu
1 5 10 15

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1 5 10 15

<210> 78
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<400> 78
Asp Trp Val Cys Glu Phe Met Lys His Gln Trp Phe Cys Asn Pro Leu
1 5 10 15

<210> 79
<211> 16
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1 5 10 15

<210> 80
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<400> 80

Asp Trp Val Cys Glu Phe Ile Lys Asn Gln Trp Met Cys Asn Val Leu
1 5 10 15

<210> 81

<211> 16

<212> PRT

<213> Artificial Sequence

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polypeptide

<400> 81

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1 5 10 15

<210> 82

<211> 16

<212> PRT

<213> Artificial Sequence

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<400> 82

Asp Trp Val Cys Glu Tyr Glu Lys Asp Gln Trp Ser Cys Asn Ile Leu
1 5 10 15

<210> 83

<211> 16

<212> PRT

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<400> 83

Asp Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Thr Leu
1 5 10 15

<210> 84

<211> 16

<212> PRT

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polypeptide

<400> 84

Asp Trp Tyr Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Val Tyr
1 5 10 15

<210> 85

<211> 16

<212> PRT

<213> Artificial Sequence

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<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 85

Asp Trp Phe Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Ser Pro Ile
1 5 10 15

<210> 86

<211> 16

<212> PRT

<213> Artificial Sequence

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polypeptide

<400> 86

Asp Trp Val Cys Glu Phe Phe Lys Lys Gln Trp Phe Cys Asn Leu Leu
1 5 10 15

<210> 87

<211> 16

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<400> 87
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1 5 10 15

<210> 88
<211> 16
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<400> 88
Asp Trp Val Cys Glu Tyr Asp Lys Gly Gln Trp His Cys Asn Ile Leu
1 5 10 15

<210> 89
<211> 16
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<400> 89
Asp Trp Ile Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Gln Gln His
1 5 10 15

<210> 90
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<400> 90

Asp Trp Val Cys Asn Trp Leu Trp Gly Gln Trp Thr Cys Asn Leu Leu
1 5 10 15

<210> 91

<211> 16

<212> PRT

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<400> 91

Asp Trp Val Cys Glu Met Phe Lys Lys Gln Trp Val Cys Asn Pro Leu
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<210> 92

<211> 16

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<400> 92

Asp Trp Ile Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Gly Pro Leu
1 5 10 15

<210> 93

<211> 16

<212> PRT

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polypeptide

<400> 93

Asp Trp Val Cys Glu Val Ile Lys Asp Gln Trp Val Cys Asn Pro Leu
1 5 10 15

<210> 94
<211> 16
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polypeptide

<400> 94
Asp Trp Val Cys Glu Asn Lys Asn Phe Lys Trp Phe Cys Asn Leu Leu
1 5 10 15

<210> 95
<211> 16
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<220>
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<400> 95
Asp Trp Val Cys Glu Tyr Ala Lys Asn Gln Trp Asn Cys Asn Pro Leu
1 5 10 15

<210> 96
<211> 16
<212> PRT
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<220>
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polypeptide

<400> 96
Asn Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Glu Trp Ala
1 5 10 15

<210> 97
<211> 16
<212> PRT
<213> Artificial Sequence

<220>

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polypeptide

<400> 97

Asn Trp Val Cys Asp Tyr Trp Lys Pro Gln Trp Phe Cys Asn Ser Tyr
1 5 10 15

<210> 98

<211> 16

<212> PRT

<213> Artificial Sequence

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polypeptide

<400> 98

Asp Trp Tyr Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Leu Val
1 5 10 15

<210> 99

<211> 16

<212> PRT

<213> Artificial Sequence

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polypeptide

<400> 99

Asn Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Glu Met
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<210> 100

<211> 16

<212> PRT

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polypeptide

<400> 100

Asp Trp Val Cys Glu Leu Phe Lys Pro Gln Trp Phe Cys Asn Ile Leu
1 5 10 15

<210> 101
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<212> PRT
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<400> 101
Asp Trp Val Cys Glu Trp Ser Lys Met Gln Trp Ser Cys Asn Ala Leu
1 5 10 15

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<400> 102
Asp Trp Val Cys Asp Tyr Lys Phe Phe Gln Trp Thr Cys Asn Leu Leu
1 5 10 15

<210> 103
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1 5 10 15

<210> 104
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<400> 104

Asp Trp Val Cys Glu Phe Phe Lys Pro Gln Trp Met Cys Asn Ile Leu
1 5 10 15

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polypeptide

<400> 105

Asp Trp Val Cys Glu Tyr Phe Lys Ser Gln Trp Met Cys Asn Met Leu
1 5 10 15

<210> 106

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<212> PRT

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<400> 106

Asp Trp Val Cys Glu Phe Phe Gly Met Gln Trp Thr Cys Asn Leu Leu
1 5 10 15

<210> 107

<211> 16

<212> PRT

<213> Artificial Sequence

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polypeptide

<400> 107

Asp Trp Val Cys Glu Tyr Ala Lys Phe Gln Trp Ile Cys Asn Ile Leu
1 5 10 15